Input Set : A:\Pto.amc

```
3 <110> APPLICANT: Brett P. Monia and Lex M. Cowsert
     5 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF TRADD EXPRESSION
     7 <130> FILE REFERENCE: RTSP-0100
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/763,748
C--> 10 <141> CURRENT FILING DATE: 2001-02-27
    12 <150> PRIOR APPLICATION NUMBER: 09/143,212
    13 <151> PRIOR FILING DATE: 1998-08-28
    15 <160> NUMBER OF SEQ ID NOS: 87
    17 <210> SEQ ID NO: 1
    18 <211> LENGTH: 1435
    19 <212> TYPE: DNA
    20 <213> ORGANISM: Homo sapiens
    22 <220> FEATURE:
    23 <221> NAME/KEY: CDS
    24 <222> LOCATION: (1)..(987)
    26 <400> SEQUENCE: 1
    27
          48
          Leu Ala Gly Val Gly Thr Gln Ala Pro Pro Arg Arg Pro Gly Gly Glu
    29
                                               10
    31
          atg gca gct ggg caa aat ggg cac gaa gag tgg gtg ggc agc gca tac
                                                                               96
          Met Ala Ala Gly Gln Asn Gly His Glu Glu Trp Val Gly Ser Ala Tyr
    32
    33
    35
          ctg ttt gtg gag tcc tcg ctg gac aag gtg gtc ctg tcg gat gcc tac
                                                                              144
          Leu Phe Val Glu Ser Ser Leu Asp Lys Val Val Leu Ser Asp Ala Tyr
    36
    37
    39
          gcg cac ccc cag cag aag gtg gca gtg tac agg gct ctg cag gct gcc
                                                                              192
    40
          Ala His Pro Gln Gln Lys Val Ala Val Tyr Arg Ala Leu Gln Ala Ala
    41
               50
    43
          ttg gca gag agc ggc ggg agc ccg gac gtg ctg cag atg ctg aag atc
                                                                              240
    44
          Leu Ala Glu Ser Gly Gly Ser Pro Asp Val Leu Gln Met Leu Lys Ile
    45
                               70
    47
          cac cgc agc gac ccg cag ctg atc gtg cag ctg cga ttc tgc ggg cgg
                                                                              288
    48
          His Arg Ser Asp Pro Gln Leu Ile Val Gln Leu Arg Phe Cys Gly Arg
    49
                                               90
          cag ccc tgt ggc cgc ttc ctc cgc gcc tac cgc gag ggg gcg ctg cgc
    51
                                                                              336
    52
          Gln Pro Cys Gly Arg Phe Leu Arg Ala Tyr Arg Glu Gly Ala Leu Arg
    53
                      100
                                          105
    55
          gee geg etg eag agg age etg geg gee geg ete gee eag eac teg gtg
                                                                              384
    56
          Ala Ala Leu Gln Arg Ser Leu Ala Ala Ala Leu Ala Gln His Ser Val
    57
                                      120
    59
          ccg ctg caa ctg gag ctg cgc gcc ggc gcc gag cqq ctq qac qct ttq
                                                                              432
    60
          Pro Leu Gln Leu Glu Leu Arg Ala Gly Ala Glu Arg Leu Asp Ala Leu
              130
    61
                                                      140
    63
          ctg gcg gac gag gag cgc tgt ttg agt tgc atc cta gcc cag cag ccc
                                                                              480
    64
          Leu Ala Asp Glu Glu Arg Cys Leu Ser Cys Ile Leu Ala Gln Gln Pro
    65
                             150
          gac cgg ctc cgg gat gaa gaa ctg gct gag ctg gag gat gcg ctg cga
    67
                                                                              528
```

Input Set : A:\Pto.amc

| 68 | Asp Arg Leu Arg Asp Glu Glu Leu Ala Glu Leu Glu Asp Ala Leu Arg | |
|------------|--|------|
| 69 | 165 170 175 | |
| 71 | aat ctg aag tgc ggc tcg ggg gcc cgg ggt ggc gac ggg gag gtc gct | 576 |
| 72 | Asn Leu Lys Cys Gly Ser Gly Ala Arg Gly Gly Asp Gly Glu Val Ala | • |
| 73 | 180 185 190 | |
| 75 76 | tog god oce ttg cag cod cog gtg cod tot otg tog gag gtg aag cog | 624 |
| 76 77 | Ser Ala Pro Leu Gln Pro Pro Val Pro Ser Leu Ser Glu Val Lys Pro | |
| 77 | 195 200 205 | 670 |
| 80 | ccg ccg ccg cca cct gcc cag act ttt ctg ttc cag ggt cag cct Pro Pro Pro Pro Pro Ala Gln Thr Phe Leu Phe Gln Gly Gln Pro | 672 |
| 81 | 210 215 220 | |
| 83 | gta gtg aat egg eeg etg age etg aag gae eaa eag aeg tte geg ege | 720 |
| 84 | Val Val Asn Arg Pro Leu Ser Leu Lys Asp Gln Gln Thr Phe Ala Arg | 720 |
| 85 | 225 230 235 240 | |
| 87 | tet gtg ggt ete aaa tgg ege aag gtg ggg ege tea etg eag ega gge | 768 |
| 88 | Ser Val Gly Leu Lys Trp Arg Lys Val Gly Arg Ser Leu Gln Arg Gly | 700 |
| 89 | 245 250 255 | |
| 91 | tgc cgg gcg ctg cgg gac ccg gcg ctg gac tcg ctg gcc tac gag tac | 816 |
| 92 | Cys Arg Ala Leu Arg Asp Pro Ala Leu Asp Ser Leu Ala Tyr Glu Tyr | 010 |
| 93 | 260 265 270 | |
| 95 | gag cgc gag gga ctg tac gag cag gcc ttc cag ctg ctg cgc ctc | 864 |
| 96 | Glu Arg Glu Gly Leu Tyr Glu Gln Ala Phe Gln Leu Leu Arg Arg Phe | |
| 97 | 275 280 285 | |
| 99 | gtg cag gcc gag ggc cgc cgc gcc acg ctg cag cgc ctg gtg gag gca | 912 |
| 100 | | |
| 101 | 290 295 300 | |
| 103 | ctc gag gag aac gag ctc acc agc ctg gca gag gac ttg ctg ggc ctg | 960 |
| 104 | Leu Glu Glu Asn Glu Leu Thr Ser Leu Ala Glu Asp Leu Leu Gly Leu | |
| 105 | 305 310 315 320 | |
| 107 | 3 3 3 3 3 3 3 3 1 3 3 3 3 3 3 3 3 3 3 3 | 1007 |
| 108 | Thr Asp Pro Asn Gly Gly Leu Ala | |
| 109 | 325 | |
| 111 | ttggagaacc tggatggcct tagggttect tetgeggeta ttgetgaacc eetgteeatc | 1067 |
| 113 | cacgggaccc tgaaactcca cttggcctat ctgctggacc tgctggggca gagttgattg | 1127 |
| 115 117 | ccttccccag gagccagacc actgggggtg catcattggg gattctgcct caggtacttt | 1187 |
| 117 | gatagagtgt ggggtggggg ggacttgctt tggagatcag cctcaccttc tcccatccca | 1247 |
| 121 | gaagegggge ttacagecag ceettacagt tteacteatg aageacettg atetttggtg | 1307 |
| 123 | tectggaett cateetgggt getgeagata etgeagtgaa gtaaaacagg aateaatett | 1367 |
| 125 | gcctgccccc agctcacact cagcgtggga ccccgaatgt taagcaatga taataaagta taacacgg | 1427 |
| | <210> SEQ ID NO: 2 | 1435 |
| | <211> LENGTH: 20 | |
| | <212> TYPE: DNA | |
| | <213> ORGANISM: Artificial Sequence | |
| | <220> FEATURE: | |
| | <223> OTHER INFORMATION: Synthetic | |
| | <400> SEQUENCE: 2 | |
| 137 | acgaggagcg ctgtttgagt | 20 |
| 1 4 0 | <210> SEQ ID NO: 3 | |

Input Set : A:\Pto.amc

| 141 | <211> LENGTH: 22 | | |
|-----|-------------------------------------|-----|----|
| | <212> TYPE: DNA | | |
| 143 | <213> ORGANISM: Artificial Sequence | | |
| 145 | <220> FEATURE: | | |
| 146 | <223> OTHER INFORMATION: Synthetic | | |
| 148 | <400> SEQUENCE: 3 | | |
| 149 | tccagctcag ccagttcttc at | | 22 |
| 152 | <210> SEQ ID NO: 4 | | |
| 153 | <211> LENGTH: 19 | | |
| 154 | <212> TYPE: DNA | • • | |
| 155 | <213> ORGANISM: Artificial Sequence | | |
| | <220> FEATURE: | | |
| 158 | <223> OTHER INFORMATION: Synthetic | , | |
| 160 | <400> SEQUENCE: 4 | | |
| 161 | ccagcagece gaceggete | | 19 |
| 164 | <210> SEQ ID NO: 5 | | |
| 165 | <211> LENGTH: 19 | | |
| 166 | <212> TYPE: DNA | | |
| 167 | <213> ORGANISM: Artificial Sequence | | |
| | <220> FEATURE: | | |
| 170 | <223> OTHER INFORMATION: Synthetic | | |
| | <400> SEQUENCE: 5 | | |
| 173 | gaaggtgaag gtcggagtc | | 19 |
| 176 | <210> SEQ ID NO: 6 | | |
| | <211> LENGTH: 20 | | |
| 178 | <212> TYPE: DNA | | |
| | <213> ORGANISM: Artificial Sequence | | |
| | <220> FEATURE: | | |
| | <223> OTHER INFORMATION: Synthetic | | |
| | <400> SEQUENCE: 6 | | |
| 185 | • | | 20 |
| 188 | <210> SEQ ID NO: 7 | | |
| | <211> LENGTH: 20 | | |
| 190 | <212> TYPE: DNA | | |
| | <213> ORGANISM: Artificial Sequence | | |
| | <220> FEATURE: | | |
| | <223> OTHER INFORMATION: Synthetic | | |
| | <400> SEQUENCE: 7 | | |
| 197 | caagetteee gtteteagee | | 20 |
| 200 | <210> SEQ ID NO: 8 | | |
| | <211> LENGTH: 18 | | |
| | <212> TYPE: DNA | | |
| | <213> ORGANISM: Artificial Sequence | | |
| | <220> FEATURE: | | |
| | <223> OTHER INFORMATION: Synthetic | | |
| | <400> SEQUENCE: 8 | | |
| 208 | ggttcccacg cccgccag | | 18 |
| | <210> SEQ ID NO: 9 | | 10 |
| | <211> LENGTH: 18 | | |
| | | | |

RAW SEQUENCE LISTING DATE: 08/03/2001 TIME: 14:04:36 PATENT APPLICATION: US/09/763,748

Input Set : A:\Pto.amc
Output Set: N:\CRF3\08032001\I763748.raw

| 213 | <212> | TYPE: DNA | | |
|-----|-------|-------------------------------|---|----|
| 214 | <213> | ORGANISM: Artificial Sequence | | •, |
| 216 | <220> | FEATURE: | | |
| 217 | <223> | OTHER INFORMATION: Synthetic | | |
| 219 | <400> | SEQUENCE: 9 | | |
| 220 | ct | cacctcct ggccgcct | | 18 |
| 223 | <210> | SEQ ID NO: 10 | | |
| 224 | <211> | LENGTH: 18 | • | |
| 225 | <212> | TYPE: DNA | | |
| 226 | <213> | ORGANISM: Artificial Sequence | | |
| | | FEATURE: | • | |
| 229 | <223> | OTHER INFORMATION: Synthetic | | |
| | | SEQUENCE: 10 | | |
| 232 | age | ctgccatc tcacctcc | · | 18 |
| 235 | | SEQ ID NO: 11 | | |
| 236 | <211> | LENGTH: 18 | | |
| 237 | <212> | TYPE: DNA | | |
| 238 | <213> | ORGANISM: Artificial Sequence | | |
| | | FEATURE: | | |
| | | OTHER INFORMATION: Synthetic | | |
| | | SEQUENCE: 11 | | |
| 244 | | cttcgtgc ccattttg | | 18 |
| 247 | | SEQ ID NO: 12 | | |
| | | LENGTH: 18 | | |
| 249 | <212> | TYPE: DNA | | |
| | | ORGANISM: Artificial Sequence | | , |
| | | FEATURE: | | |
| | | OTHER INFORMATION: Synthetic | | |
| | | SEQUENCE: 12 | | |
| 256 | | cccactct tcgtgccc | | 18 |
| 259 | | SEQ ID NO: 13 | • | |
| | | LENGTH: 18 | • | |
| 261 | <212> | TYPE: DNA | | |
| | | ORGANISM: Artificial Sequence | | |
| | | FEATURE: | | |
| | | OTHER INFORMATION: Synthetic | | |
| | | SEQUENCE: 13 | | |
| 268 | | geceace caetette | | 18 |
| | _ | SEQ ID NO: 14 | | |
| | | LENGTH: 18 | | |
| | | TYPE: DNA | | |
| | | ORGANISM: Artificial Sequence | | |
| | | FEATURE: | | |
| 277 | <223> | OTHER INFORMATION: Synthetic | | |
| | | SEQUENCE: 14 | | |
| 280 | | atgeget geceacee | | 18 |
| | | SEQ ID NO: 15 | | |
| | | LENGTH: 18 | | |
| | | TYPE: DNA | | |
| | | | | |

Input Set : A:\Pto.amc

| | 286 | <213> | ORGANISM: Artificial Sequence | |
|---|-----|-------|---|----|
| : | 288 | <220> | FEATURE: | |
| : | 289 | <223> | OTHER INFORMATION: Synthetic | |
| | | | SEQUENCE: 15 | |
| | | | cacaaaca ggtatgcg | 18 |
| | | | SEQ ID NO: 16 | |
| | | | LENGTH: 18 | |
| | | | TYPE: DNA | |
| | | | ORGANISM: Artificial Sequence FEATURE: | |
| | | | OTHER INFORMATION: Synthetic | |
| | | | SEQUENCE: 16 | |
| | 304 | | ggactcca caaacagg | 18 |
| | | | SEQ ID NO: 17 | 10 |
| | | | LENGTH: 18 | |
| | | | TYPE: DNA | |
| | | | ORGANISM: Artificial Sequence | |
| | | | FEATURE: | |
| | | | OTHER INFORMATION: Synthetic | |
| | | | SEQUENCE: 17 | |
| ; | 316 | gto | ccagcgag gactccac | 18 |
| 3 | 319 | <210> | SEQ ID NO: 18 | |
| | 320 | <211> | LENGTH: 18 | |
| | | | TYPE: DNA | |
| 3 | 322 | <213> | ORGANISM: Artificial Sequence | |
| | | | FEATURE: | |
| | | | OTHER INFORMATION: Synthetic | |
| | | | SEQUENCE: 18 | |
| | | | ecttgtc cagcgagg | 18 |
| | | | SEQ ID NO: 19 | |
| | | | LENGTH: 18 | |
| | | | TYPE: DNA | |
| | | | ORGANISM: Artificial Sequence | |
| | | | FEATURE: | |
| | | | OTHER INFORMATION: Synthetic SEQUENCE: 19 | |
| | 340 | | _ | 18 |
| | | _ | teegaca ggaceace SEQ ID NO: 20 | 10 |
| | | | LENGTH: 18 | |
| | | | TYPE: DNA | |
| | | | ORGANISM: Artificial Sequence | |
| | | | FEATURE: | |
| | | | OTHER INFORMATION: Synthetic | |
| | | | SEQUENCE: 20 | |
| | 52 | | • | 18 |
| 3 | 55 | | SEQ ID NO: 21 | |
| | | | LENGTH: 18 | |
| 3 | 57 | <212> | TYPE: DNA | |
| 3 | 58 | <213> | ORGANISM: Artificial Sequence | |
| | | | | |

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/763,748

DATE: 08/03/2001 TIME: 14:04:37

Input Set : A:\Pto.amc

Output Set: N:\CRF3\08032001\1763748.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:207 M:283 W: Missing Blank Line separator, <400> field identifier